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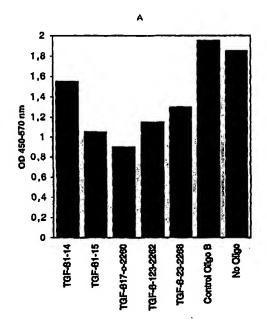
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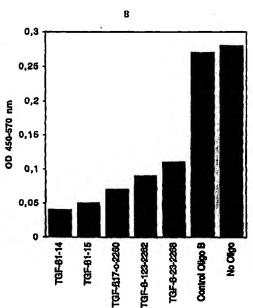
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(54) Title: A METHOD FOR STIMULATING THE IMMUNE SYSTEM





(57) Abstract

Medicament comprising a combination of at least one inhibitor of the effect of a substance negatively effecting an immune response, the substance selected from the group consisting of $TGF-\beta$ and its receptors, VEGF and its receptors, interleukin 10 (IL-10) and its receptors, PGE₂ and its receptors, wherein the inhibitor has a molecular weight of less than 100 kDa and at least one stimulator positively effecting an immune response.

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A Method for Stimulating the immune system

Two different approaches have been used in the prior art to enhance the immune response against neoplastic cells. One approach uses the addition of cytokines like interleukin-2 (IL-2) or transfection of tumor cells and/or immune cells with genes coding for cytokines like IL-2 or other proteins enhancing the immune response like transfection of tumor cells with lymphotactin or like transfection of T-lymphocytes with CD-40 Ligand.

The second approach uses the inhibition of immunosuppressive molecules to enhance the body's immune response to tumor cells. Thus, J. NEUROSURG. 78 (1993) 944-51, Jachimczak et al. (1993) and WO 94/25588, Schlingensiepen et al. (1994) teach the use of antisense oligonucleotides targeted to TGF-\(\beta\) to reverse tumor-induced immunosuppression.

Several documents in the prior art teach that a combination of these two approaches is either not efficacious or is not beneficial over use of one of the two approaches used alone.

Thus, CANCER BIOTHER. 8(2), 1993, 159 - 170, Gridley et al., as well as CANCER BIOTHER. 9(4), 1994, 317-327, Mao et al., both teach that a combination of anti-transforming growth factor-beta antibody with IL-2 does not cause significant antitumor effects.

Furthermore, PROC. NATL. ACAD. SCI 93, (1996), 2909-2914, Fakhrai et al., teaches that a combination of transfection with genes encoding antisense sequences to transforming growth factor beta (TGF-\$\beta\$) TGF-\$\beta\$ mRNA with transfection of IL-2 into tumor cells does not increase the immune response against the tumor compared to transfection with TGF-\$\beta\$ antisense alone.

Surprisingly, in contrast, certain combinations of stimulators and inhibitors are more efficacious than either approach alone.

The present invention discloses a medicament comprising a combination of

- at least one inhibitor of the effect of a substance negatively effecting an immune response, the substance selected from the group consisting of TGF-B and its receptors, VEGF and its receptors, interleukin 10 (IL-10) and its receptors, PGE₂ and its receptors, wherein the inhibitor has a molecular weight of less than 100 kDa and
- at least one stimulator positively effecting an immune response.

In a preferred embodiment, the inhibitor is inhibiting the synthesis or function of molecules suppressing or downregulating or negatively affecting the immune response. The inhibitor can be an oligonucleotide which may function as an antisense nucleotide or a ribozyme or it may be an antibody fragment derived from an antibody e.g. a fab-fragment or a single chain antibody.

Preferably, the stimulator is positively effecting the immune response by increasing presentation of antigens and/or enhancing proliferation and/or function of immune cells.

In a preferred embodiment, the stimulator is enhancing the synthesis or function of molecules stimulating, enhancing, upregulating and/or positively regulating the immune response. In particular, the stimulator is stimulating and/or enhancing the synthesis and/or the function of factors such as GM-CSF, SCF, CSF, IFN-γ, FLT-3-ligand as well as monocyte chemotatic proteins (MCP-1), interleukin-2, interleukin-4, interleukin-12 and/or interleukin-18 or is one of the mentioned interleukins or is selected from the group consisting of viruses, viral antigens, antigens expressed in tumor cells or pathogens but not in normal cells, organspecific antigenes expressed in affected organs which are not essential for the organism, e. g. prostate, ovary, breast, melanine producing cells.

The stimulators are preferably selected from

- a) Chemokines, including lymphotactin and/or immune cell attracting substances and/or
- b) viruses and/or parts of viruses, including retroviruses, adenoviruses, papillomaviruses, Epstein-Barr-Vriuses, Viruses that are non-pathogenic including Newcastle-Disease virus, Cow-pox-virus and/or
- c) autologous and/or heterologous MHC-Molecules and/or
- d) molecules involved in antigen processing and/or
- e) molecules involved in antigen presentation and/or
- f) molecules involved in mediating immune cell effects and/or
- g) molecules involved in mediating immune cell cytotoxic effects and/or

- h) molecules involved in antigen transportation and/or
- i) co-stimulatory molecules
- j) peptides enhancing recognition by immune cells and/or cytotoxic effects of immune cells
- k) the peptides containing one or more amino acids differing between a protein in the target cell from the other cells within an organism
- 1) the peptides according to j) being
 - Peptides containing one or more mutations and/or amino acid substitutions of the ras protein amino and/or
 - Peptides containing one or more mutations and/or amino acid substitutions of the p53 protein and/or
 - Peptides containing one or more mutations and/or amino acid substitutions of the EGF-Receptor protein and/or
 - Peptides containing one or more mutations and/or amino acid substitutions of fusion peptides and/or fusion proteins and/or
 - Peptides containing one or more mutations and/or amino acid substitutions and/or amino acid substitutions caused by gene rearrangements and/or gene translocations and/or
 - Peptides containing one or more mutations and/or amino acid substitutions of the retinoblastoma protein and/or

- Peptides containing one or more mutations and/or amino acid substitutions of proteins coded by oncogenes and/or protooncogenes and/or
- Peptides containing one or more mutations and/or amino acid substitutions of proteins coded by anti-oncogenes and/or tumor suppressor genes and/or
- Peptides derived from proteins differing in the target cell by one or more amino acids from the proteins expressed by other cells in the same organism and/or
- Peptides derived from viral antigens and/or coded by viral nucleic acids and/or
- Peptides derived from proteins expressed in a diseased organ but not in the nervous system, muscle, hematopoetic system or other organs essential for survival. Diseased organs are e. g. prostate, ovary, breast, melanine producing cells and the like.
- m) tumor cell extracts and/or tumor cell lysates and/or adjuvants,
- n) fusion cells of dendritic and tumor cells.

These fusion cells are hybridoma cells derived from a mixture of dentritic cells and tumor cells. Dentritic cells are generated e. g. by treatment of PBMC with GM-CSF and IL-4 or a mixture of GM-CSF, IL-4 and IFN-γ or FLT-3 ligand. Fusion of dendritic cells with tumor cells can be achieved e. g. using PEG (polyethylene glycole) or electrofusion.

Surprisingly, treatment of PBMC with VEGF-oligonucleotides enhanced the number and/or effectiveness of dendritic cells.

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In one embodiment of the invention the inhibitor is an oligonucleotide. Preferably the oligonucleotides of Fig. 1 are useful in the medicament of the present invention.

In a further embodiment, the invention provides oligonucleotides having one of the sequences given in figure 1-2 to 1-4.

Also oligonucleotides having 1 to 10 additional nucleotides at the 5'- or 3'- end are part of the invention.

Oligonucleotide sequences used for transfection are usually much longer sequences than those used for antisense oligonucleotides, which usually do not exceed 30 bases in length and are applied as short single-stranded sequences and are not integrated into a vector system.

Since transfected sequences are usually much longer than oligonucleotides, if cross inhibition of different members of a protein family would occur with the antisense technology, such cross inhibition of other mRNAs than the target mRNA, is much more likely with transfected antisense sequences, compared to oligonucleotides. However, Cell Growth Differ, Vol. 6(12), February 1995, pages 1635 - 1642, Huang, F. et al.teaches "only the K6 transfectant exhibited 39 and 33% respectively of the levels or TGF beta1 mRNA and active secreted TGF beta1 protein of the parental line. K6 exhibited no change in TGF beta2 expression and TGF beta3 expression was not detected in either parental or transfectant cell line."

It was therefore surprising to find oligonucleotides according to this invention, which were able to significantly reduce expression of both, TGF- β_1 as well as TGF- β_2 e. g. TGF- β_1 -14, TGF- β_1 -15, TGF- β_2 -17-c-2260, TGF- β_2 -123-2262, TGF- β_2 -23-2268, TGF- β_2 -4, TGF- β_2 -14, TGF- β_2 -15, TGF- β_2 -9, TGF- β_2 -14/1, TGF- β_2 -14/2, TGF- β_1 -136. Furthermore surprisingly oligonucleotides were designed, which were

able to significantly reduce expression of TGF-\(\beta_2\) as well as TGF-\(\beta_3\).

Surprisingly even oligonucleotides were found, which were able to significantly reduce expression of TGF- β_2 as well as TGF- β_1 and TGF- β_3 , e. g. b1-N17, b1-N14, b1-N24, TGF- β_2 -9, TGF- β_2 -14, TGF- β_2 -215, TGF- β_3 -17-c-2260, TGF- β_3 -12-9/20-2261, TGF- β_3 -123-2262, TGF- β_3 -12-9/22-2263, TGF- β_3 -23-2268, TGF- β_3 -98-11, TGF- β_3 -98-23, TGF- β_3 -98-7, TGF- β_3 -98-10, TGF- β_3 -1-rwk-5, TGF- β_3 -3-rwk-2, TGF- β_3 -1-rwk-5, TGF- β_3 -3-rwk-2, TGF- β_3 -1-rwk-5, TGF- β_3 -1-rwk-10.

Thus oligonucleotides which are effective against expression of at least two of TGF- β_1 , TGF- β_2 and/or TGF- β_3 are also part of the invention.

These findings were also surprising in view of the fact that sequence comparison between the mRNAs of TGF- β_2 , TGF- β_1 and TGF- β_3 showed that not a single sequence of 20 bases in length could be found that would be identical within the three different mRNAs. Even if such a hypothetical sequence had really existed, inhibition of the three mRNAs by such a hypothetical consensus sequence would have been extremely unlikely, since it is well known in the art that only a small minority of antisense sequences complementary to a certain mRNA actually exert a so-called antisense effect, *i.e.* inhibit expression of the respective protein.

Endothelial synthesis of monocyte chemotactic protein-1 (MCP-1) has been implicated in the regulation of monocyte recruitment for extravascular pools both under physiological and inflammatory conditions.

MCP-1 antisense oligonuclotides were able to modulate monocyte infiltration and were thus anti-inflammatoric.

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These antisense-oligonucloetides are useful for the treatment of inflammatory diseases e.g. asthma, morbus crohn, collitis ulcerosa, diabetes, glomerulonephritis, acute respiratory distress syndrome and artherosclerotic plaque formation.

In a preferred embodiment of the invention the oligonucleotides and/or ribozymes and/or nucleic acids have modifications at the bases, the sugars and/or the phosphate moieties of the oligonucleotides.

In a further preferred embodiment of the invention the oligonucleotides and/or ribozymes and/or nucleic acids have modifications wherein the modifications are phosphorothicate (S-ODN) internucleotide linkages and/or methylphosphonate internucleotide linkages and/or phosphoramidate linkages and/or peptide linkages and/or 2'-O-derivatives, such as 2'-O-methyl or 2'-O-methoxyethoxy modifications of the sugar and/or modifications of the bases.

In a further preferred embodiment of the invention the oligonucleotides and/or ribozymes and/or nucleic acids are coupled to or mixed with folic acid, hormones, steroid hormones such as oestrogene, progesterone, corticosteroids, mineral corticoids, peptides, proteoglycans, glycolipids, phospholipids, polyethylene imine or other poly cations and derivatives therefrom.

Furthermore, the present invention provides a method of treating hyperproliferative diseases, neoplasms or infectious diseases by administering a medicament of the invention to patients in need thereof. The method is especially useful for the treatment of leukemia, non-hodgkin lymphoma, hodgkin lymphoma, bronchial carcinoma, esophageal carcinoma, colorectal carcinoma, gastric carcinomas, intestinal tumors, hepatic tumors, gall bladder and gallduct carcinomas, pancreatic carcinoma, anal carcinoma, breast cancer, ovarian carcinoma, cervial carcinoma, endometrium carcinoma, prostatic carcinoma, bladder carcinoma, malignant melanoma, brain tumors, and sarcomas.

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The necessary doses of the medicament of the present invention depend on the disease and the severity of the disease. Whereas higher levels are more effective, they often have a higher degree of side effects. Suitable doses are selected to obtain concentrations of the oligonucleotides in the range of 0.1 to 10 µmol/l and concentrations of the cytokines in the range of 10 to 1.000 U/ml in the patient blood.

In a preferred embodiment of the invention the inhibitor of the effect of a substance negatively effecting an immune response is applied locally to a tumor or other pathologically affected site or organ and the stimulator positively effecting an immune response is applied systemically (e.g. i.v. or s.c. or orally).

In another preferred embodiment of the invention the inhibitor of the effect of a substance negatively effecting an immune response is applied systemically (e.g. i.v or s.c. or orally) to the tumor and the stimulator positively effecting an immune response is applied locally to a tumor or other pathologically affected site or organ. In another preferred embodiment of the invention the inhibitor of the effect of a substance negatively effecting an immune response is applied systemically (e.g. i.v. or s.c. or orally) to the tumor and the stimulator positively effecting an immune response is applied systemically (e.g. i.v. or s.c. or orally).

In another preferred embodiment of the invention the inhibitor of the effect of a substance negatively effecting an immune response is applied locally to a tumor or other pathologically affected site or organ and the stimulator positively effecting an immune response is applied locally to a tumor or other pathologically affected site or organ.

Fig. 1 shows oligonucleotides useful in the present invention.

1.

Fig. 2A shows effects of oligonucleotides (f.c. 5 μ M) on TGF- β 2 secretion in glioma cells in 10% MEM Dulbecco medium (3 day incubation with oligonucleotides).

Fig. 2B shows effects of oligonucleotides (f.c. 5 μM) on TGF-β1 secretion in PBMC in 10% FCS RPMI 1640 medium (3 day incubation with oligonucleotides).

Fig. 3A shows effects of oligonucleotides (f.c. 5 μ M) on TGF-B1 secretion in PBMC in 10% FCS RPMI 1640 medium (3 day incubation with oligonucleotides).

Fig. 3B shows effects of oligonucleotides (f.c. 5 μM) on TGF-β2 secretion in glioma cells in 10% FCS RPMI 1640 medium (3 day incubation with oligonucleotides).

Fig. 4A shows TGF-B1 concentration (ELISA) in glioma cells (3 day incubation with oligonucleotides).

Fig. 4B shows TGF-ß2 concentration (ELISA) in glioma cells (3 day incubation with oligonucleotides).

Fig. 5 shows lysis of tumor-cells: LAK-Cytotoxicty, Ratio of glioma-cells/PBMC: 1:20.

Fig 6A shows dendritic cells generated from PBMC (% of control). Cytokines: GM-CSF (400 U/ml) + IL-4 (300 U/ml).

Fig. 6B shows lysis of tumor-cells: Effects of 5 μM VEGF-Antisense-Oligos on LAK-Cytotoxicty. Ration of tumor-cells/DC/PBMC was 1:5:20.

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Fig. 7A shows effects of oligonucleotides (f.c. 5 μM) on TGF-β1 secretion in PBMC in 10% FCS RPMI 1640 medium (3 day incubation with oligonucleotides).

Fig. 7B shows effects of oligonucleotides (f.c. 5 μ M) on TGF- β 2 secretion in tumro cells in 10% FCS RPMI 1640 medium (3 day incubation with oligonucleotides).

Fig. 8 shows lysis of tumor-cells: Effects of oligonucleotides on LAK-Cytotoxicty. Ration of tumor-cells/PBMC was 1:20.

Examples

Preparation of PBMC and tumor cells

Peripheral blood mononuclear cells (PBMC) were isolated from venous blood of healthy donors by standard Ficoll-Hypaque gradient centrifugation. Briefly, heparinized blood was mixed with equal volumes of complete medium (CM: RMPI 1640 medium supplemented with 10% (v/v) fetal calf serum and 1 mM L-Glutamine) and layered onto a Ficoll-Hypaque (Pharmacia, Uppsala, Sweden) gradient. After centrifugation at 400g for 30 min at room temperature, PBMCs banded at the plasma-Ficoll interface were recovered, washed tree times and resuspended in complete medium. Cell viability, as determined by Trypan blue exclusion, was greater than 97%.

Human glioma cell lines were established from tumor specimens of patients with anaplastic astrocytoma (WHO Grad III) or from glioblastoma (WHO Grad IV).

Measurement of cell proliferation

For PBMC-proliferation assays (3H-thymidine incorporation and cell counting), freshly isolated PBMCs were cultured for 72h in 96-well round-bottom plates (Nunc, Copenhagen, Denmark) at a final concentration (f.c.) of 10⁵ cells/well (100 µl CM). For cell number determination the cells were counted by hemacytometer. Cell viability was determined by trypan blue staining. Treated and untreated cells showed 95-100% viability after 72h in vitro growth (with or without S-ODN).

For the tumor proliferation experiments 10⁴/100 µL glioma cells were seeded into 96-well flat-bottom plates (Nunc, Denmark) and incubated with cytokines and/or oligonucleotides. The DNA synthesis rate was measured by a standard 3H-thymidine incorporation assay and determination of cell number was performed as described above.

Quantification of TGF-\$1 protein in culture supernatants by enzyme-linked immunosorbent assay (ELISA)

The culture medium was harvested after 3 days, cleared of cellular components by centrifugation, filtered and stored at -70°C until processed further. TGF-\(\beta\)1 and TGF-\(\beta\)2 concentrations were measured after acidification of supernatants by TGF-\(\beta\)1 and TGF-\(\beta\)2 ELISA (R&D Systems, Minneapolis, USA) in duplicates, as recommended by the manufacturer.

Figures 1 - 4 and 7 show the effect of oligonucleotides on the TGF-ß secretion in cells. The concentration of the TGF-ß is reported as an optical density. The higher the optical density the higher is the concentration of the TGF-ß.

Figure 1A and 1B shows the effect of the oligonucleotides on the TGF-ß secretion.

Control oligos (GAA GGA ATT ACC ACT TTC) have no effects whereas the oli-

gonucleotides shown in the figures reduce the secretion of TGF-\(\beta\). The oligos in figure 1 are more effective against TGF-\(\beta\)1.

Figure 2 shows further oligos and their effects on TGF-ß secrection. TGF-ß-14 is especially effective against the secrection of TGF-ß1 and -ß2.

Figure 3 shows further oligonucleotides being effective against secretion of TGF-\(\beta\)1 and -\(\beta\)2. These oligonucleotides are more effective against TGF-\(\beta\)2 but are also effective against TGF-\(\beta\)1.

Figure 8 shows a supra additive effect on tumor cell cytotoxicity by a combination of 2 μ M each of a TGF-B1 and TGF-B2 antisense oligonucleotide compared to a single 5 μ M dose of either oligonucleotide.

CARE-LASS (calcein-release-assay) to measure cytotoxic PBMC activity

A standard calcein-release-assay (CARE-LASS assay) to determine cytotoxic activity of PBMC was employed as described by Lichtenfels, R., Biddison, W.E., Schulz, H., Vogt, A.B. and R. Martin. CARE-LASS (calcein-release assay), an improved fluorescence-based test system to measure cytotoxic lymphocyte activity. J. Immunol. Meth., 172: 227-239, 1994.

Target and Effector cells

At the day of the assay malignant glioma were harvested, washed twice in 5% FCS /PBS and incubated with Calcein-AM (Molecular Probes, USA) for 30min in 37°C. Labeled target cells were washed twice in 5% FCS/PBS, adjusted to 100 000 / ml, and plated into 96-well U-shaped microtiter plates (Nunc, Dennmark) at the final volumen of 100uL/well.

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PBMC were washed with 5% FCS/PBS and adjusted to final concentration of 1-10 Mio cells /ml.

Cells were treated with cytokines and oligodeoxynucleotides as described in the individual experiments.

Assay

To measure CTL activity effector cells were plated into 96-well U-shape microtiter plates at Target: Effector Ratios of 1:10 - 1:100. To measure spontanous release and total release of calcein, wells were preloaded with 200uL 5%FCS/PBS or 200uL lysis buffer (50mM sodium-borate, 0.1% Triton, pH 9.0) respectively. After incubating the plate for 4 h at 37°C in an incubator, 100uL of supernatans were transferred into new wells and measured employing an automated fluorescence scanner (Titertek Fluoroskan II, Germany). Both for excitation and for emission, filter settings 2 were chosen (ex 2 – 485nm, em 2 –538 nm). The percent of cytotoxicity was determined from the following equation:

F/CTL assay - F spontaneous release
----- x 100 = % cytotoxicity
F total lysis - F spontanous release

In one set of experiments, glioma cells, denritic cells (DC) and PBMC were cocultured. In these experiments DC were generated from PBMC using the cytokines GM-CSF and IL4. Cells were further treated with antisense VEGF-oligonucleotides according to the invention or with no oligonucleotides as control experiments. Tumor cells were also treated with the cytokines GM-CSF and IL4 with or without oligonudeotides.

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PBMC were only treated with oligonucleotides according to the invention, but not with the cytokines GM-CSF and IL4. oligos were used at a concentration of 5 μ M unless indicated otherwise in the descriptions in the figures.

The CARE-LASS (calcein-release-assay) was used to measure cytotoxic PBMC activity.

In one set of experiments glioma cells and PBMC were treated either with a single oligonucleotide or with a combination of oligonucleotides. The single oligonucleotides were given at 5 μ M concentration. In the combination experiment, each oligonucleotide was given at 2 μ M concentration. Both, PBMC and tumor cells were incubated separately with the oligonucleotide(s) for 72 h.

The CARE-LASS (calcein-release-assay) was used to measure cytotoxic PBMC activity.

Claims

- 1. Medicament comprising a combination of
 - at least one inhibitor of the effect of a substance negatively effecting an immune response, the substance selected from the group consisting of TGF-B and its receptors, VEGF and its receptors, interleukin 10 (IL-10) and its receptors, PGE₂ and its receptors, wherein the inhibitor has a molecular weight of less than 100 kDa and
 - at least one stimulator positively effecting an immune response.
- 2. The medicament of claim 1 wherein the inhibitor is inhibiting the synthesis or function of molecules suppressing or downregulating or negatively affecting the immune response.
- 3. The medicament of claim 1 wherein the inhibitor is an oligonucleotide.
- 4. The medicament according to claim 3 wherein the oligonucleotide is an antisense nucleotide and/or a ribozyme.
- 5. The medicament according to claim 3 wherein the oligonucleotides has a sequence according to figure 1.
- 6. The medicament according to claim 1, wherein the inhibitor is a fab-fragment or single chain antibody (scFv).
- 7. The medicament according to claim 1, wherein the stimulator is enhancing the synthesis or function of molecules stimulating, enhancing, upregulating and/or positively regulating the immune response.

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- 8. The medicament according to claim 7, wherein the stimulator is stimulating and/or enhancing the synthesis and/or the function of factors such as GM-CSF, SCF, CSF, IFN, FLT-3-ligand, monocyte chemotatic proteins (MCP-1), interleukin-2, interleukin-4, interleukin-12 and/or interleukin-18 or is one of the mentioned interleukins or is selected from the groups consisting of viruses, viral antigens, antigens expressed in tumor cells or pathogens, but not in normal cells, organ specific antigens expressed in affected organs which are not essential for the organism or fusion cell of dendritic and tumor cells.
- 9. The medicament according to claim 1, wherein the medicament comprises two or more of the inhibitors and/or the stimulators.
- An oligonucleotide having one of the sequences of figures 1-2 to 1-4 (No. 55 -213).
- 11. The oligonucleotide according to claim 10 wherein each oligonucleotide is effective against expression of at least two of TGF-β₁, TGF-β₂ and/or TGF-β₃ having the sequence.
- 12. A method of treating neoplasm or infectious diseases by administering a medicament according to claim 1 to a patient in need thereof.
- 13. A method according to claim 12 for the treatment of hyperproliferative diseases, leukemia, non-hodgkin lymphoma, hodgkin lymphoma, bronchial carcinoma, esophageal carcinoma, colorectal carcinoma, gastric carcinomas, intestinal tumors, hepatic tumors, gall bladder and gallduct carcinomas, pancreatic carcinoma, anal carcinoma, mastocarcinoma, ovarian carcinoma, cervial carcinoma, endometrium carcinoma, prostatic carcinoma, bladder carcinoma, malignant melanoma, brain tumors, and/or sarcomas.

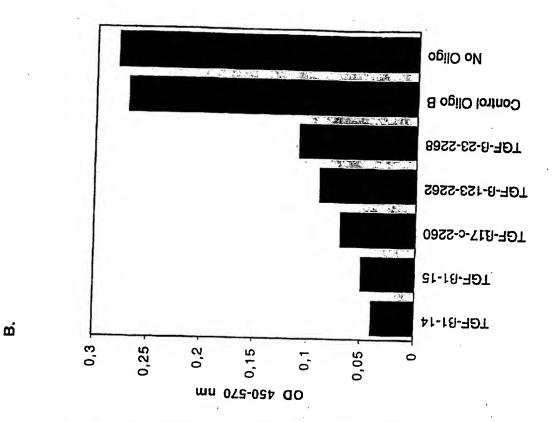
1.	TGF-B2-1	C ACA CAG TAG TGC A
2.	TGF-B2-2	GC ACA CAG TAG TGC
3.	TGF-82-3	GC TTG CTC AGG ATC TGC
4.		
	TGF-B2-4	TAC TCT TCG TCG CT
5.	TGF-B2-5	C TTG GCG TAG TAC T
6.	TGF-B2-6	G TAA ACC TCC TTG G
7.	TGF-B2-7	GT CTA TTT TGT AAA CCT CC
8.	TGF-B2-8	GC ATG TCT ATT TTG TAA ACC
9.	TGF-B2-9	CGG CAT GTC TAT TTT GTA
10.	TGF-B2-10	G GCA TCA AGG TAC C
11.	TGF-82-11	CTG TAG AAA GTG GG
12.	TGF-B2-12	AC AAT TCT GAA GTA GGG T
13.	TGF-B2-13	T CAC CAA ATT GGA AGC AT
14.	TGF-B2-14	GCT TTC ACC AAA TTG GAA GC
15.	TGF-B2-15	CTG GCT TTT GGG TT
16.	TGF-B2-16	T CTG ATA TAG CTC AAT CC
17.	TGF-B2-17	T CCT AGT GGA CTT TAT AG
18.	TGF-B2-18	T TTT TCC TAG TGG ACT
19.	TGF-B2-19	C AAT TAT CCT GCA CAT TTC
20.	TGF-82-20	GC AAT TAT CCT GCA CA
21.	TGF-B2-21	GC AGC AAT TAT CCT GC
22.	TGF-B2-22	TG GCA TTG TAC CCT
23.	TGF-B2-23	TG TGC TGA GTG TCT
24.	TGF-B2-24	CC TGC TGT GCT GAG TG
25.	TGF-82-25	C TTG GGT GTT TTG C
26.	TGF-B2-26	T TTA GCT GCA TTT GCA AG
27.	TGF-B2-27	G CCA CTT TTC CAA G
28.	TGF-B2-14/1	CTT TCA CCA AAT TGG AAG
29.	TGF-B2-14/2	CAC CAA ATT GGA AGC
30.	TGF-B2-14/3	TCA CCA AAT TGG AAG C
31.	TGF-82-15/1	CTC TGG CTT TTG GG
		CGG CAT GTC TAT TTT G
32.	TGF-B2-9/1	CGG CAI GIC IAI III G
33.	መሮፑ 01 1	ככא שאר שכש שכר אר
	TGF-81-1	CGA TAG TCT TGC AG
34.	TGF-B1-2	GTC GAT AGT CTT GC
35.	TGF-B1-3	CTT GGA CAG GAT CT
36.	TGF-B1-4	CCA GGA ATT GTT GC
37.	TGF-B1-5	CCT CAA TTT CCC CT
38.	TGF-B1-6	GAT GTC CAC TTG CA
39.	TGF-B1-7	CTC CAA ATG TAG GG
40.	TGF-B1-8	ACC TTG CTG TAC TG
41.	TGF-B1-9	GTA GTA CAC GAT GG
42.	TGF-81-10	CAC GTA GTA CAC GA
43	TGF-B1-11	CAT GTT GGA CAG CT
44.	TGF-81-12	GCA CGA TCA TGT TG
45.	TGF-B1-13	TGT ACT CTG CTT GAA C
46.	TGF-81-14	CTG ATG TGT TGA AGA ACA
47.	TGF-B1-15	CTC TGA TGT GTT GAA G
48.	TGF-B1-16	GGA AGT CAA TGT ACA G
49.	TGF-B1-17	CAT GTC GAT AGT CTT GCA
50.	TGF-B1-18	AGC TGA AGC AAT AGT TGG
51.	TGF-B1-19	GTC ATA GAT TTC GTT GTG
52.	TGF-81-20	CTC CAC TTT TAA CTT GAG
53.	TGF-81-21	TGC TGT ATT TCT GGT ACA
54.	TGF-B1-137	CGA TAG TCT TGC AG
J 1 .	101 111-111	CO.1 1110 101 100 MG

1 / 11 Fig. 1-1

55.	b1-N17	TCC TCT TCG ACT GCT CTC	
56.	b2-N14	CGA AGG TTA AAC CAC TTT CG	
57.	b2-N24	GTG AGT CGT GTC GTC C	
58.	TGF-B2-98-1	CATCGTTGTCGTCG	
59.	TGF-B2-98-2	CGCTTCTTCCGCCG	
60.	TGF-B2-98-3	CGAAGGAGCCATTCG	
61.	TGF-B2-98-4	CGATGTAGCG	
62.	TGF-B2-98-5	CGTCAAATCG	
63.	TGF-B2-98-6	CGTAGTACTCTTCGTCG	
64.	TGF-B2-98-7	CGCGCTCGCAGGCG	•
65.	TGF-B2-98-8	CGGCCGCCTCCGGCTCG	
66.	TGF-B2-98-9	CGCGGATCGCCTCG	
67.	TGF-B2-98-10	GAGCGCGACCGTGAC	
07.	1Gr -132 - 38 - 10	GAGCGCGACCGIGAC	
68.	TGF-B-17-c-2260	ACC TCC TTG GCG TAG TA	
69.	TGF-B-12-9/20-2261	AGG GCG GCA TGT CTA TTT TG	
70.	TGF-B-123-2262	CAG AAG TTG GCA TTG TAC	
	TGF-B-12-9/22-2263		
71.	· · · · · ·	AGG GCG GCA TGT CTA TTT TGT A	
72.	TGF-B-23-2268	TGG GAC ACG CAG CAA GG	
73.	TGF-81-98-1	CGGGGGCGGGGG	
74.	TGF-81-98-2	CGGGGGGGGGGGG	
75.	TGF-\$1-98-3	CGGCGCCGAGGCGCCCG	
76.	TGF-B1-98-4	CCGAGGTCCTTGCGG	
77.	TGF-B1-98-5	CGGCGGTGCCGGGA	
78.	TGF-B1-98-6	CTCGGCGGCCGGTAG	
79.	TGF-81-98-7	CGCTAAGGCG	
80.	TGF-81-98-8	CCGCACAACTCCGG	
81.	TGF-81-98-9	GCGAGTCGCTGG	
82.		CGGTTGCTGAGGTATCG	
83.	TGF-81-98-11	CCGGGAGAGCAACACGG	
84.	TGF-B1-98-12	CGCTTCTCG	
85.	TGF-81-98-13	CCATTAGCACGCGG	
86.	TGF-B1-98-14	CGGGCTCCG	
87.	TGF-B1-98-15	CCGGCCACCCGGTCGCGG	
88.	TGF-B1-98-16	CGAGCACGGCCTCG	
89.	TGF-B1-98-17	CGGGCAGCGGGCCG	
90.	TGF-B1-98-18	CGCGGATGGCCTCG	
91.	TGF-B1-98-19	CGATGCGCTTCCG	
92.	TGF-B1-98-20	CCCGCGGCGGG	
93.	TGF-B1-98-21	CGCAGCCCGGAGGGCG	
94.	TGF-B1-98-22	CGGCGCCCCCG	
95.	TGF-B1-98-23	CGGCACTGCCGAGAGCGCG	
96.	TGF-B1-98-24	CGGGGATGAAGGCGGCG	
97.	TGF-B1-98-25	CGGGTCGGCGACTCCCG	
98.	TGF-B1-98-26	CGCCTGAGGGACGCCG	
99.	TGF-81-98-27	AAGCGTCCCGGCG	•
100.	TGF-81-98-28	CGCGGGCAGCGTCGCG	
101.	TGF-81-98-29	CCCCGCGCCTCCGG	
102.	TGF-81-98-30	CGGCGGCTCG	
103.	TGF-B1-98-31	CGCTCCGGGCCGAGGCCG	
104.	TGF-81-98-32	CGGCCCGCGGGCG	
105.	TGF-81-98-33	CGGACGGGCGTCC	
106.	TGF-\$1-98-34	CGGCCGGGCCCTCG	Fig. 1-2
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	107.	TGF-B3-98-1	TCGAGCTTCCCCGA
	108.	TGF-B3-98-2	CCCGGAGCCGAAGG
	109.	TGF-B3-98-3	CCCGAGGAGCGGG
	110.	TGF-B3-98-4	ACGCAGCAAGGCGA
	111.	TGF-B3-98-5	CGGGTTGTCGAGCCG
	112.	TGF-B3-98-6	CGGCAGTGCCCCG
	113.	TGF-B3-98-7	CGGAATTCTGCTCG
	114.	TGF-B3-98-8	TTCGTTGTGCTCCG
	115.	TGF-ß3-98-9	ATTCCGACTCGGTG
	116.	TGF-B3-98-10	ACGTGGGTCATCACCGT
	117.	TGF-B3-98-11	CGAAGAAGCG
	118.	TGF-ß3-312	CCT AAT GGC TTC CA
	119.	VEGF-98-1	CGGCCGCGTGTGT
	120.	VEGF-98-2	CGGGAATGCTTCCGCCG
	121.	VEGF-98-3	CGGCTCACCGCCTCGGC
	122.	VEGF-98-4	CACGTCTGCGGATC
	123.	VEGF-98-5	CCCCGCATCGCATCAGGG
	124.	VEGF-98-6	CGCCTTGCAACGCG
	125.	VEGF-98-7	CCGACCGGGCCGG
	126.	VEGF-49	GTTCATGGTTTCGG
	127.	VEGF-55	GCAGAAAGTTCATGG
	128.	VEGF-188	GCTGATAGACATCC
	129.	VEGF-190	GCGCTGATAGACAT
	130.	VEGF-194	GTAGCTGCGCTGATAG
	131.	VEGF-253	CTCGATCTCATCAG
		VEGF-255	ATGTACTCGATCTCATC
	133.	VEGF-260	GAAGATGTACTCGATC
		VEGF-263	CTTGAAGATGTACTCG
	135.	VEGF-292	GCATCGCATCAGGG
	136.	VEGF-294	CCGCATCGCATCAG
	137.	VEGF-422	CATTTGTTGTGCTGTAGG
	138.	VEGF-434	GGTCTGCATTCACATTTG
	139.	VEGF-441	CTTTGGTCTGCATTC
	140.	VEGF-445	CTTTCTTTGGTCTGC
	141.	VEGF-450	GCTCTATCTTTGG
		VEGF-455	GTCTTGCTCTATCTTTC
	143.	VEGF-459	CTTGTCTTGCTCTATC
	144.	VEGF-596	CATCTGCAAGTACGTTCG
	145.	VEGF-598	CACATCTGCAAGTACGTT
		VEGF-600	GTCACATCTGCAAGTACG
		VEGF-600-2	CATCTGCAAGTACG
		VEGF-601	CACATCTGCAAGTAC
	149.	VEGF-604	GTCACATCTGCAAG
	150.	VEGF-607	CTTGTCACATCTGC
	151.	VEGF-607-2	GGCTTGTCACATCTGC
	152.	VEGF-610	CTCGGCTTGTCACATC
	153.	VEGF-638	CTCCTTCCTGC
	154.	VEGF-766	GCT TGA AGA TGT ACCT CG
	155.	VEGF-r-1062	CGT TGC TCT CCG ACG
	156.	flt-1165	GAC ACG GCC TTT TCG
٠,	157.	flt-rm-2115	CCA GCA GCT GAC CAT GG
	158.	flk1/kdr-m-2315	GAA ATC GAC CCT CGG
	159.	MCP-1-Rec-A/B-571	GCA TGT TGT GGA TG
	160.	MCP-1-1954	GCA GAG ACT TTC ATG C
	161.	MCP-1-1955	ATA ACA GCA GGT GAC TGG Fig. 1-3
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			2 / 11

		,
162.	MCP-1-1956	GAA CCC ACT TCT GC
163.	MCP-1-2761	GAC ACT TGC TGC TG
164.	MCP-1-2762	CCA CTT CTG CTT GGG
165.	VEGF-703	CTG CAA GTA CGT TCG
	flt-1567	TCC CTT ATG ATG CCA GCA AGT G
167.		CCA GCA ATG ACA GC
168.		G GGA AAG CTG AGG C
169.		T CGA GGG AAA GCT GA
	TGF-ß-1-rwk-3	C CTC GAG GGA AAG C
	TGF-B-1-rwk-4	GG GCT GGT GTG
	TGF-B-1-rwk-5	
		GA ACA GGG CTG GTG TG
	TGF-ß-1-rwk-6	G AAC AGG GCT GGT G
	TGF-ß-1-rwk-7	AG AGC GCG AAC AGG
	TGF-ß-l-rwk-8	GA GAG CGC GAA CAG G
176.		CGA GAG CGC GAA CAG
	TGF-B-1-rwk-10	CCC CTG GCT CGG GGG
	TGF-B-1-rwk-11	C CCT GGC TCG GGG
	TGF-B-1-rwk-12	C CCC TGG CTC GGG G
	TGF-B-1-rwk-13	TCC CCC TGG CTC GG
181.		C TCC CCC TGG CTC G
182.	TGF-B-1-rwk-15	TGC GCT TCC GCT TCA C
183.	TGF-ß-l-rwk-16	CC TCG ATG CGC TTC
184.	TGF-ß-l-rwk-17	G ATG GCC TCG ATG C
185.	TGF-ß-1-rwk-18	G GAT GGC CTC GAT GC
186.	TGF-ß-l-rwk-19	ATG GCC TCG ATG CGC TT
187.	TGF-ß-3-rwk-1	TC AGC AGG GCC AGG
188.	TGF-B-3-rwk-2	GCA AAG TTC AGC AGG GC
189.	TGF-ß-3-rwk-3	GG CAA AGT TCA GCA GG
190.	TGF-B-3-rwk-4	GT GGC AAA GTT CAG CAG G
191.		GTG GCA AAG TTC AG
192.		GAC CGT GGC AAA GTT CAG
193.	TGF-ß-3-rwk-7	AGA GAG GCT GAC CGT
194.	TGF-B-3-rwk-8	GAC AGA GAG AGG CTG AC
195.	TGF-ß-3-rwk-9	A CAG AGA GAG GCT GA
196.	TGF-B-3-rwk-10	GT GGA CAG AGA GAG G
197.	TGF-8-3-rwk-11	CA AGT GGA CAG AGA GAG G
198.	TGF-B-3-rwk-12	TCT TCT TGA TGT GGC C
199.	TGF-8-3-rwk-13	CC CTC TTC TTC TTG ATG
200.	TGF-B-3-rwk-14	C ACC CTC TTC TTC T
201.	TGF-B-3-rwk-15	A TGG ATT TCT TTG GCA T
202.	TGF-ß-3-rwk-16	GGA TTT CTT TGG C
203.	TGF-B-3-rwk-17	AA GTT GGA CTC TCT TCT C
204.	TGF-B-3-rwk-18	TAA GTT GGA CTC TCT TCT
205.	TGF-B-3-rwk-19	GAC CTA AGT TGG ACT C
206.	TGF-B-3-rwk-20	T TTC TAG ACC TAA GTT GG
207.	TGF-B-3-rwk-21	CT GAT TTC TAG ACC TAA G
208.	TGF-8-3-rwk-22	G AAG CAG TAA TTG GTG T
209.	TGF-B-3-rwk-23	GG AAT CAT CAT GAG G
210.	TGF-8-3-rwk-24	GGG AAT CAT CAT GAG
211.	TGF-B-3-rwk-25	G GTT GTC GAG CCG GT
212.	TGF-8-3-rwk-26	GTC CTC CCA ACA TAG TA
213.	TGF-B-3-rwk-27	GG GTC CTC CCA ACA
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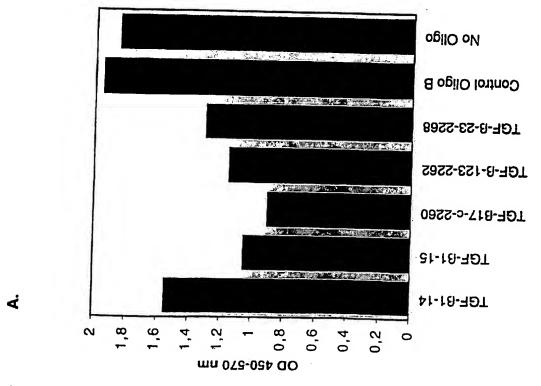


Figure 2

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PCT/EP99/04013

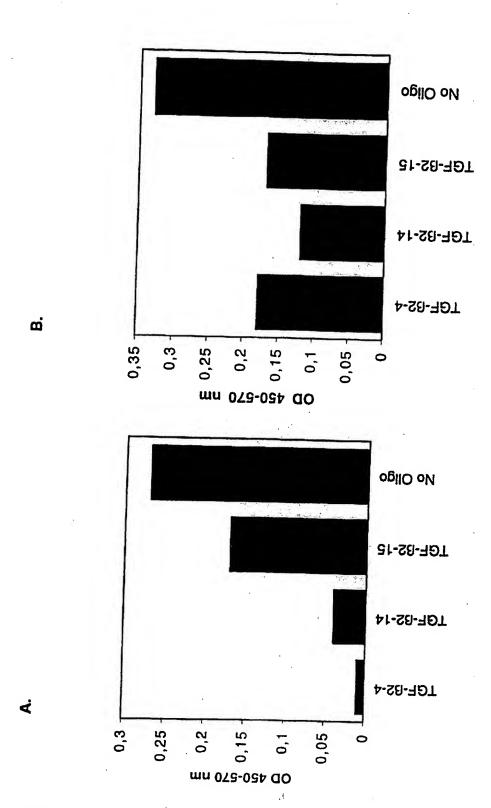
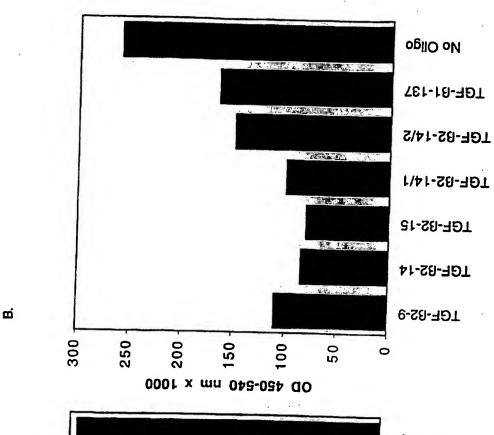


Figure 3

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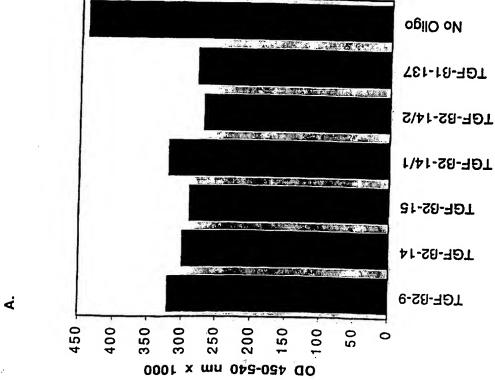


Figure 4

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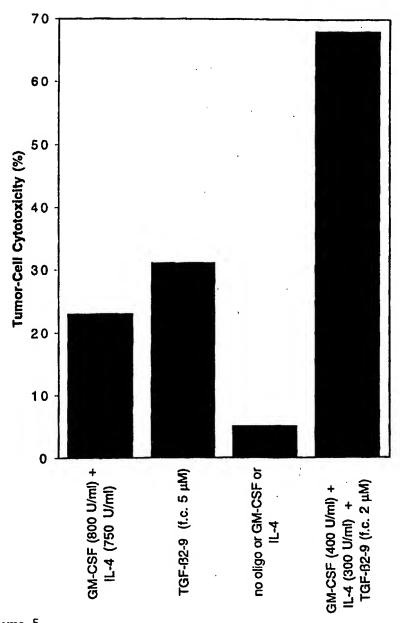
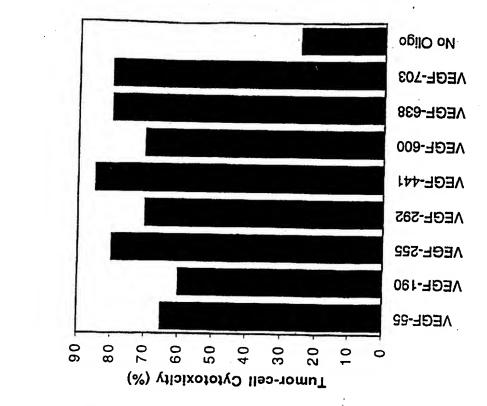
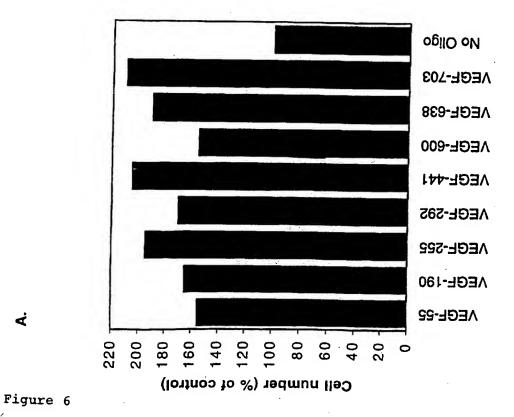


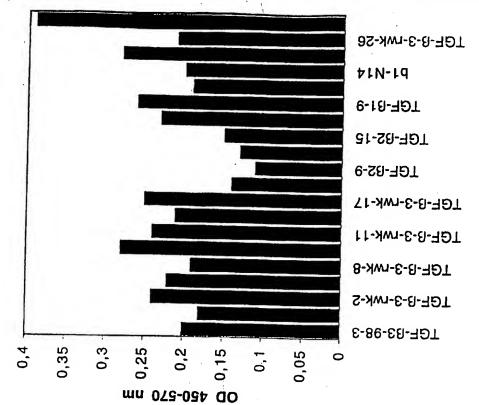
Figure 5

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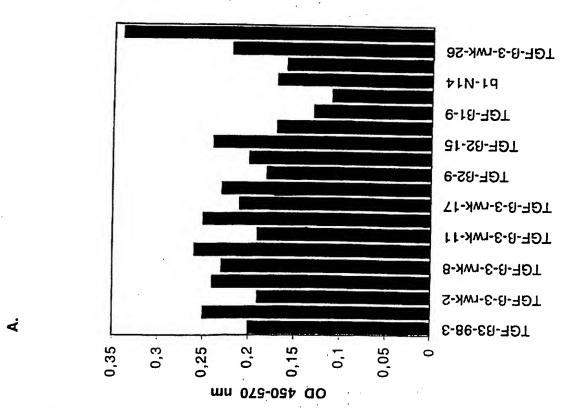


Figure 7

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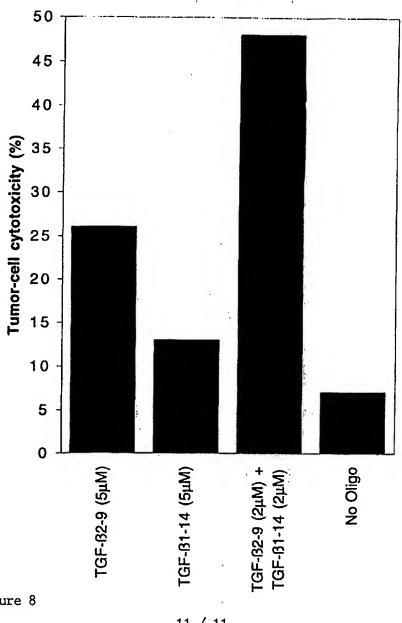


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